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Application Serial Number:

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
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 Room 1B03, Arlington, VA 22202

Revised 05/17/04



RAW SEQUENCE LISTING

DATE: 10/04/2004

PATENT APPLICATION: US/10/509,248

TIME: 11:15:48

30

Input Set : A:\PTO.FG.txt .

Output Set: N:\CRF4\10042004\J509248.raw

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3 <110> APPLICANT: Japan Science and Technology Corporation
         5 <120> TITLE OF INVENTION: Growth factors ior the likej presented protein hollow nano-
   particles for
                 therapeutic drug .
        7 <130> FILE REFERENCE: P023P02
   C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/509,248
        9 <141> CURRENT FILING DATE: 2004-09-28
         9 <150> PRIOR APPLICATION NUMBER: JP 2002-97395
                                                                        Does Not Comply
   W--> 10 <151> PRIOR FILING DATE: 2002-3-29
                                                                        Corrected Diskette Needed
        12 <160> NUMBER OF SEQ ID NOS: 4
        14 <170> SOFTWARE: PatentIn Ver. 2.1
        16 <210> SEQ ID NO: 1
        17 <211> LENGTH: 39
        18 <212> TYPE: DNA
        19 <213> ORGANISM: (Artificial Sequence
     -> 20 (<220> FEATURE) Synthesized Oligonucleotide
     -> 1 <223> (OTHER INFORMATION:)
       21 <400> SEQUENCE: 1
        22 cgacaaggca tgggaggcgg ccgcagccct caggctcag
   24 <210> SEQ ID NO: 2
       25 <211> LENGTH: 39
        26 <212> TYPE: DNA
      27 <213> ORGANISM Artificial Sequence
    --> 28 <220> FEATURE: Synthesized Oligonucleotide
   W--> 29 (223) OTHER INFORMATION:
     -> 29 <400> SEQUENCE: 2
DIANK 30 ctgagcctga gggctgcggc cgcctcccat gccttgtcg
   all 32 <210> SEQ ID NO: 3
      33 <211> LENGTH: 27
   nes. 34 <212> Type: DNA
        35 <213> ORGANISM Artificial Sequence
   W--> 36 (220) FEATURE Synthesized Oligonucleotide
   W--> 37 <223 OTHER INFORMATION:
   W--> 37 <400> SEQUENCE: 3
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40 <210> SEQ ID NO: 4 PAME (220> blowk
        41 <211> LENGTH: 30
        42 <212> TYPE: DNA
   43 <213> ORGANISM: Artificial Sequence
W--> 44 <220> FEATURE: Synthesized Oligonucleotide
   W--> 45 <223> OTHER INFORMATION:
   W--> 45 <400> SEQUENCE: 4
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46 gggcggccgc cacgcagttc ccaccatttc

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/509,248

DATE: 10/04/2004 TIME: 11:15:49

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\10042004\J509248.raw

Use of <220> Feature(NEW RULES):

From Explanation on

gequence(s) are missing the <220> Feature and associated headings.

Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence"

or"Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104,pp.29631-32)

(Sec.1.823 of new Rules)

Seq#:1,2,3,4

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/509,248

DATE: 10/04/2004 TIME: 11:15:49

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\10042004\J509248.raw

L:7 M:283 W: Missing Blank Line separator, <130> field identifier L:9 M:270 C: Current Application Number differs, Replaced Current Application No L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:10 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD

L:20 M:283 W: Missing Blank Line separator, <220> field identifier L:20 M:256 W: Invalid Numeric Header Field, <220> has non-blank data

L:21 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:1, <213> ORGANISM: Artificial Sequence

L:21 M:283 W: Missing Blank Line separator, <400> field identifier

L:21 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:21

L:28 M:283 W: Missing Blank Line separator, <220> field identifier

L:28 M:256 W: Invalid Numeric Header Field, <220> has non-blank data

L:29 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:2, <213> ORGANISM: Artificial Sequence

L:29 M:283 W: Missing Blank Line separator, <400> field identifier

L:29 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:2,Line#:29

L:36 M:283 W: Missing Blank Line separator, <220> field identifier

L:36 M:256 W: Invalid Numeric Header Field, <220> has non-blank data

L:37 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:3, <213> ORGANISM: Artificial Sequence

L:37 M:283 W: Missing Blank Line separator, <400> field identifier

L:37 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:37

L:44 M:283 W: Missing Blank Line separator, <220> field identifier L:44 M:256 W: Invalid Numeric Header Field, <220> has non-blank data

L:45 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:4, <213> ORGANISM: Artificial Sequence

L:45 M:283 W: Missing Blank Line separator, <400> field identifier

L:45 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:4,Line#:45